## IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

New Atty. Dock t No: 37629-0004 Old Atty. Docket No. 047744/0106

Group Art Unit: 1631

Examiner: M. Zeman

re patent application of RE( NAPPIK, Achim et al.

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Application No.: 09/490, PECH CENTER 1600/2900

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**AMENDMENT** 

Commissioner for Patents Washington, D.C. 20231

Sir:

TECH CENTER 1600/2900 In response to the Office Action mailed May 7, 2002, Applicants respectfully request that the following amendments be entered into the application:

## IN THE SPECIFICATION:

## Page 18, line 4, paragraph 2: Please amend as follows:

Figs. 2A-2G: Alignment of consensus sequences designed for each subgroup (amino acid residues are shown with their standard one-letter abbreviation). (2A-2B) (SEQ ID NOS 28-31, respectively) kappa sequences, (2C-2D) (SEQ ID NOS 32-34, respectively) lambda sequences and (2E-2G) (SEQ ID NOS 35-41, respectively), heavy chain sequences. The positions are numbered according to Kabat (1991). In order to maximize homology in the alignment, gaps (-) have been introduced in the sequence at certain positions.

## Page 18, line 10, paragraph 3: Please amend as follows:

Figs. 3A-3K: Gene sequences (SEQ ID NOS 42, 44, 46 and 48, respectively) of the synthetic V kappa consensus genes. The corresponding amino acid sequences (SEQ ID NOS 43, 45, 47 and 49, respectively) (see Figs. 2A-2B) as well as the unique cleavage sites are also shown.